

O/S/0 04-18-01 O/PE #2

4/5/2001

Serial Number: 09/813,718

CRF Processing Date:

Edited by:

Verified by:

(STIC Staff)

 Changed a file from non-ASCII to ASCII **ENTERED** Changed the margins in cases where the sequence text was wrapped down to the next line. Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was  the prior application data; or  other \_\_\_\_\_ Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted:  non-ASCII "garbage" at the beginning/end of files;  secretary initials/filename at end of file;  
 page numbers throughout text;  other invalid text, such as \_\_\_\_\_ Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: Other:

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/19/95

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/813,718

DATE: 04/11/2001

TIME: 17:38:24

Input Set : A:\pto.da

Output Set: N:\CRF3\04112001\I813718.raw

3 <110> APPLICANT: Schimmel, Paul  
 4 Wakasugi, Keisuke  
 6 <120> TITLE OF INVENTION: Human Aminoacyl-tRNA Synthetase Polypeptides Useful For  
 7 The Regulation of Angiogenesis  
 9 <130> FILE REFERENCE: 00-221  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/813,718  
 C--> 12 <141> CURRENT FILING DATE: 2001-03-21  
 14 <160> NUMBER OF SEQ ID NOS: 58  
 16 <170> SOFTWARE: PatentIn Ver. 2.0  
 18 <210> SEQ ID NO: 1  
 19 <211> LENGTH: 5174  
 20 <212> TYPE: DNA  
 21 <213> ORGANISM: Artificial Sequence  
 23 <220> FEATURE:  
 24 <221> NAME/KEY: CDS  
 25 <222> LOCATION: (3428)..(5035)  
 27 <220> FEATURE:  
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 29 full-length TyrrRS in pET20B  
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 34 cagcgtgacc gctacacttg ccagccctt agcgcccgct ctttcgcctt tcttccttc 120  
 36 ctttctcgcc acgttcgcgg gctttcccg toaagotota aatcgggggc tccctttagg 180  
 38 gttccgattt agtgctttac ggcacctcgaa ccccaaaaaa cttgatagg gtgatggttc 240  
 40 acgttgtgg ccatcgccctt gatagacggt ttttcgcctt ttgacgttgg agtccacgtt 300  
 42 cttaatagt ggactcttgtt tccaaactgg aacaacactc aaccatatct cggcttattc 360  
 44 ttttgattttt taagggattt tgccgatttc ggcctattgg taaaaaaaaatg agctgattta 420  
 46 aaaaaatttt aacgcgaattt ttaacaaaat attaacgtttt acaatttcag gtggcacttt 480  
 48 tcggggaaat gtgcgcggaa cccctatttg ttatatttttcaaaatacatt caaatatgt 540  
 50 tccgctcatg agacaataaac cctgataaaat gttcaataaa tattgaaaaaa ggaagagtat 600  
 52 gagtattcaa catttccgtt tgcgccttat tccctttttt gcggcattttt gccttcctgt 660  
 54 ttttgctcac ccagaaacgc tggtaaaatg aaaagatgctt gaagatcgtt tgggtgcacg 720  
 56 agtgggttac atcgaacttgg atctcaacag cggttaagatc cttgagatgtt ttgcggccg 780  
 58 aagaacgtttt ccaatgtatgaa gcactttaa agttctgttata tttggcgcgg tattatccc 840  
 60 tattgacgcc gggcaagagc aactcggctcg cccgcatacac tatttctcaga atgacttgg 900  
 62 tgagtactca ccagtcacag aaaagatctt tacggatggc atgacagtaa gagaattatg 960  
 64 cagtgctgcc ataaccatgtt gtgataacac tgcggccaac ttacttctgtt caacgatcg 1020  
 66 aggaccgaag gagctaaccg ctttttgcataacatgggg gatcatgttata ctcgccttgc 1080  
 68 tcgttggaa ccggagctgtt atgaagccat accaaacgcgac gacgtgaca ccacgatgcc 1140  
 70 tgcagcaatg gcaacaacgt tgcgcacactt attaactggc gaactactt ctctagctt 1200  
 72 ccggcaacaa ttaatagact ggatggaggc ggataaaatgtt gcaggaccac ttctgcgtt 1260  
 74 gcccctccg gctggctgtt ttatgttgcataa atctgttgcataa gcccgttgcgttgcgtt 1320  
 76 cggatccatg gcaagacttgg ggccagatgg taagccctcc cgtatctgtt ttatctacac 1380  
 78 gacggggagt caggcaactt tggatgttgcataa atatagacag atcgctgaga taggtgcctc 1440  
 80 actgatgttgcataa ctttgcataa agtttactca tatatactttt agattgttgcataa 1500  
 82 aaaacttcat ttttaatttta aaaggatctt ggtgaagatc cttttgcataa atctcatgac 1560  
 84 caaaatccctt taacgtgagt tttcggttccat ctgagcgttca gacccgttcaaaaatccctt 1620

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PATENT APPLICATION: US/09/813,718

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Input Set : A:\Pto.da

Output Set: N:\CRF3\04112001\I813718.raw

86	aggatttct	tgagatccott	tttttctgcg	cgtaatctgc	tgcttgcaaa	caaaaaacc	1680										
88	accgctacca	gcgggtgggtt	gttgcggga	tcaagagcta	ccaactctt	ttccgaaggt	1740										
90	aactggcttc	agcagagcgc	agataccaaa	tactgtcctt	ctagtgttagc	cgtagttagg	1800										
92	ccaccacttc	aagaactctg	tagcaccggc	tacataccctc	getctgtcaa	tcctgttacc	1860										
94	agtggctgt	gccagtgccg	ataagtctgt	tcttaccggg	ttggactcaa	gacgatagtt	1920										
96	accggataag	gcgcagcgg	cgggctgaac	ggggggttcg	tgcacacagc	ccagcttgg	1980										
98	gcgaacgacc	tacaccgaac	tgagataacct	acagcgttag	ctatgagaaa	gcgccacgct	2040										
100	tcccgaaggg	agaaaggccg	acaggtatcc	ggtaagcgc	agggtcggaa	caggagagcg	2100										
102	cacgagggag	cttccagggg	gaaacgcctg	gtatcttat	agtccctgtcg	ggttgcucca	2160										
104	cctctgactt	gagcgtcgat	ttttgtatg	ctcgtcaggg	gggcggagcc	tatggaaaaa	2220										
106	cggcagcaac	gcggcctttt	tacggttcct	ggccttttgc	tggccttttgc	ctcacatgtt	2280										
108	cttccctgct	ttatccccgt	attctgtgga	taaccgtatt	accgccttgc	agtgagctga	2340										
110	taccgctcgc	cgcagccaa	cgaccggagcg	cagcgagtc	gtgaggcgagg	aagcggaaaga	2400										
112	gcccgtatg	cggatttttc	tccttacgca	tctgtcggt	atttcacacc	gcatatatgg	2460										
114	tgcactctca	gtacaatctg	ctctgatgccc	gcatagttaa	gccagttatac	actccgtat	2520										
116	cgtacgtga	ctgggtcatg	gctgcgcccc	gacacccgccc	aacacccgct	gacgcgcct	2580										
118	gacgggcttg	tctgctcccg	gcatccgctt	acagacaaga	tgtgaccgctc	tccggagct	2640										
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122	catcagcgtg	gtcgtgaago	gattcacaga	tgtctgcctg	ttcattccgc	tccagctcg	2760										
124	tga <sup>g</sup> tttctc	cagaagcgtt	aatgtctggc	ttctgtataaa	gcggccatg	ttaagggcgg	2820										
126	tttttctgt	tttggtaact	gatgcctccg	tgttaagggg	atttctgttc	atggggtaa	2880										
128	tgataccgat	gaaacgagag	aggatgctca	cgatacggt	tactgtatgt	gaacatgc	2940										
130	ggttactgga	acgttgtgag	ggtaaacaaac	tggcggtatg	gatcggcg	gaccagagaa	3000										
132	aaatc <sup>c</sup> ctca	gggtcaatgc	cagcgcttc	ttaatacaga	tgttaggttt	ccacagggt	3060										
134	gccagcagca	tcctgcgtat	cagatccgga	acataatgtt	gcagggcgt	gacttccgc	3120										
136	tttccagact	ttacgaaaca	cgaaaccga	agaccattca	tgttgttgc	caggcgcag	3180										
138	acgttttgc	gcagcagtc	cttcacgttc	gctcgctat	cgggtattca	ttctgtcaac	3240										
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142	cccggtggca	ggacccaacg	ctgcccggaga	tctcgatccc	gcgaattaa	tacgactcac	3360										
144	tatagggaga	ccacaacgg	ttccctctag	aaataatttt	gtttaacttt	aagaaggaga	3420										
146	tatacat atg	ggg gac	gtt ccc	agc ctt	gaa gag	aaa ctg	cac ctt atc	3469									
147	Met	Gly	Asp	Ala	Pro	Ser	Pro	Glu	Glu	Lys	Leu	His	Leu	Ile			
148	1	5	10														
150	acc	cg	aa	ctg	cag	gag	gtt	ctg	ggg	gaa	gag	aag	ctg	aag	gag	ata	3517
151	Thr	Arg	Asn	Leu	Gln	Glu	Val	Leu	Gly	Glu	Glu	Lys	Leu	Lys	Glu	Ile	
152	15				20				25							30	
154	ctg	aag	gag	cgg	gaa	ctt	aaa	att	tac	tgg	gga	acg	gca	acc	acg	ggc	3565
155	Leu	Lys	Glu	Arg	Glu	Leu	Lys	Ile	Tyr	Trp	Gly	Thr	Ala	Thr	Thr	Gly	
156									35							45	
158	aaa	cca	cat	gtg	gtc	tac	ttt	gtg	ccc	atg	tca	aag	att	gca	gac	ttc	3613
159	Lys	Pro	His	Val	Ala	Tyr	Phe	Val	Pro	Met	Ser	Lys	Ile	Ala	Asp	Phe	
160									50							60	
162	tta	aag	gca	ggg	tgt	gag	gtt	aca	att	ctg	ttt	gct	gac	ctc	cac	gca	3661
163	Leu	Lys	Ala	Gly	Cys	Glu	Val	Thr	Ile	Leu	Phe	Ala	Asp	Leu	His	Ala	
164									65							75	
166	tac	ctg	gat	aac	atg	aaa	gcc	cca	tgg	gaa	ctt	cta	gaa	ctc	cga	gtc	3709
167	Tyr	Leu	Asp	Asn	Met	Lys	Ala	Pro	Trp	Glu	Leu	Leu	Glu	Leu	Arg	Val	
168									80							90	
170	agt	tac	tat	gag	aat	gtt	atc	aaa	gca	atg	ctg	gag	agc	att	ggt	gtt	3757

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/813,718

DATE: 04/11/2001  
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Output Set: N:\CRF3\04112001\I813718.raw

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174	ccc	ttg	gag	aag	ctc	aag	ttc	atc	aaa	ggc	act	gat	tac	cag	ctc	agc	3805
175	Pro	Leu	Glu	Lys	Leu	Lys	Phe	Ile	Lys	Gly	Thr	Asp	Tyr	Gln	Leu	Ser	
176					115				120							125	
178	aaa	gag	tac	aca	cta	gat	gtg	tac	aga	ctc	tcc	tcc	gtg	gtc	aca	cag	3853
179	Lys	Glu	Tyr	Thr	Leu	Asp	Val	Tyr	Arg	Leu	Ser	Ser	Val	Val	Thr	Gln	
180					130				135							140	
182	cac	gat	tcc	aag	gct	gga	gct	gag	gtg	gta	aag	cag	gtg	gag	cac		3901
183	His	Asp	Ser	Lys	Lys	Ala	Gly	Ala	Glu	Val	Val	Lys	Gln	Val	Glu	His	
184					145				150							155	
186	cct	ttg	ctg	agt	ggc	ctc	tta	tac	ccc	gga	ctg	cag	gct	ttg	gat	gaa	3949
187	Pro	Leu	Leu	Ser	Gly	Leu	Leu	Tyr	Pro	Gly	Leu	Gln	Ala	Leu	Asp	Glu	
188					160				165							170	
190	gag	tat	tta	aaa	gta	gat	gcc	caa	ttt	gga	ggc	att	gat	cag	aga	aag	3997
191	Glu	Tyr	Leu	Lys	Val	Asp	Ala	Gln	Phe	Gly	Gly	Ile	Asp	Gln	Arg	Lys	
192	175				180				185							190	
194	att	ttc	acc	ttt	gca	gag	aag	tac	ctc	cct	gca	ctt	ggc	tat	tca	aaa	4045
195	Ile	Phe	Thr	Phe	Ala	Glu	Lys	Tyr	Leu	Pro	Ala	Leu	Gly	Tyr	Ser	Lys	
196					195				200							205	
198	cgg	gtc	cat	ctg	atg	aat	cct	atg	gtt	cca	gga	tta	aca	ggc	agc	aaa	4093
199	Arg	Val	His	Leu	Met	Asn	Pro	Met	Val	Pro	Gly	Leu	Thr	Gly	Ser	Lys	
200					210				215							220	
202	atg	agc	tct	tca	gaa	gag	gag	tcc	aag	att	gat	ctc	ctt	gat	cgg	aag	4141
203	Met	Ser	Ser	Ser	Glu	Glu	Glu	Ser	Lys	Ile	Asp	Leu	Leu	Asp	Arg	Lys	
204					225				230							235	
206	gag	gat	gtg	aag	aaa	aaa	ctg	aag	aag	gcc	ttc	tgt	gag	cca	gga	aat	4189
207	Glu	Asp	Val	Lys	Lys	Lys	Leu	Lys	Lys	Ala	Phe	Cys	Glu	Pro	Gly	Asn	
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210	gtg	gag	aac	aat	ggg	gtt	ctg	tcc	ttc	atc	aag	cat	gtc	ctt	ttt	ccc	4237
211	Val	Glu	Asn	Asn	Gly	Val	Leu	Ser	Phe	Ile	Lys	His	Val	Leu	Phe	Pro	
212	255				260				265							270	
214	ctt	aag	tcc	gag	ttt	gtg	atc	cta	cga	gat	gag	aaa	tgg	ggg	gga	aac	4285
215	Leu	Lys	Ser	Glu	Phe	Val	Ile	Leu	Arg	Asp	Glu	Lys	Trp	Gly	Gly	Asn	
216					275				280							285	
218	aaa	acc	tac	aca	gct	tac	gtg	gac	ctg	gaa	aag	gac	ttt	gct	gct	gag	4333
219	Lys	Thr	Tyr	Thr	Ala	Tyr	Val	Asp	Leu	Glu	Lys	Asp	Phe	Ala	Ala	Glu	
220					290				295							300	
222	gtt	gta	cat	cct	gga	gac	ctg	aag	aat	tct	gtt	gaa	gtc	gca	ctg	aac	4381
223	Val	Val	His	Pro	Gly	Asp	Leu	Lys	Asn	Ser	Val	Glu	Val	Ala	Leu	Asn	
224					305				310							315	
226	aag	ttg	ctg	gat	cca	atc	cgg	gaa	aag	ttt	aat	acc	cct	gcc	ctg	aaa	4429
227	Lys	Leu	Leu	Asp	Pro	Ile	Arg	Glu	Lys	Phe	Asn	Thr	Pro	Ala	Leu	Lys	
228					320				325							330	
230	aaa	ctg	gcc	agc	gct	gcc	tac	cca	gat	ccc	tca	aag	cag	aag	cca	atg	4477
231	Lys	Leu	Ala	Ser	Ala	Ala	Tyr	Pro	Asp	Pro	Ser	Lys	Gln	Lys	Pro	Met	
232	335				340				345							350	
234	gcc	aaa	ggc	cct	gcc	aag	aat	tca	gaa	cca	gag	gag	gtc	atc	cca	tcc	4525
235	Ala	Lys	Gly	Pro	Ala	Lys	Asn	Ser	Glu	Pro	Glu	Glu	Val	Ile	Pro	Ser	

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Input Set : A:\Pto.da  
Output Set: N:\CRF3\04112001\I813718.raw

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239	Arg Leu Asp Ile Arg Val Gly Lys Ile Ile Thr Val Glu Lys His Pro			
240	370	375	380	
242	gat gca gac agc ctg tat gta gag aag att gac gtg ggg gaa gct gaa			4621
243	Asp Ala Asp Ser Leu Tyr Val Glu Lys Ile Asp Val Gly Glu Ala Glu			
244	385	390	395	
246	cca cgg act gtg gtg agc ggc ctg gta cag ttc gtg ccc aag gag gaa			4669
247	Pro Arg Thr Val Val Ser Gly Leu Val Gln Phe Val Pro Lys Glu Glu			
248	400	405	410	
250	ctg cag gac agg ctg gta gtg gtg ctg tgc aac ctg aaa ccc cag aag			4717
251	Leu Gln Asp Arg Leu Val Val Leu Cys Asn Leu Lys Pro Gln Lys			
252	415	420	425	430
254	atg aga gga gtc gag tcc caa ggc atg ctt ctg tgt gct tct ata gaa			4765
255	Met Arg Gly Val Ser Gln Gly Met Leu Leu Cys Ala Ser Ile Glu			
256	435	440	445	
258	ggg ata aac cgc cag gtt gaa cct ctg gac cct ccg gca ggc tct gct			4813
259	Gly Ile Asn Arg Gln Val Glu Pro Leu Asp Pro Pro Ala Gly Ser Ala			
260	450	455	460	
262	cct ggt gag cac gtg ttt gtg aag ggc tat gaa aag ggc caa cca gat			4861
263	Pro Gly Glu His Val Phe Val Lys Gly Tyr Glu Lys Gly Gln Pro Asp			
264	465	470	475	
266	gag gag ctc aag ccc aag aag aaa gtc ttc gag aag ttg cag gct gac			4909
267	Glu Glu Leu Lys Pro Lys Lys Val Phe Glu Lys Leu Gln Ala Asp			
268	480	485	490	
270	tcc aaa att tct gag gag tgc atc gca cag tgg aag caa acc aac ttc			4957
271	Phe Lys Ile Ser Glu Glu Cys Ile Ala Gln Trp Lys Gln Thr Asn Phe			
272	495	500	505	510
274	atg acc aag ctg ggc tcc att tcc tgt aaa tcg ctg aaa ggg ggg aac			5005
275	Met Thr Lys Leu Gly Ser Ile Ser Cys Lys Ser Leu Lys Gly Gly Asn			
276	515	520	525	
278	att agc ctc gag cac cac cac cac tgagatccgg ctgctaaca			5055
279	Ile Ser Leu Glu His His His His His			
280	530	535		
282	agcccgaaag gaagctgagt tggctgctgc caccgcttag caataactag cataaccct	5115		
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288	<211> LENGTH: 536			
289	<212> TYPE: PRT			
290	<213> ORGANISM: Artificial Sequence			
292	<220> FEATURE:			
293	<223> OTHER INFORMATION: Description of Artificial Sequence: human			
294	full-length TyrRS in pET20B			
297	<400> SEQUENCE: 2			
298	Met Gly Asp Ala Pro Ser Pro Glu Glu Lys Leu His Leu Ile Thr Arg			
299	1	5	10	15
301	Asn Leu Gln Glu Val Leu Gly Glu Lys Leu Lys Glu Ile Leu Lys			
302	20	25	30	
304	Glu Arg Glu Leu Lys Ile Tyr Trp Gly Thr Ala Thr Thr Gly Lys Pro			

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305	35	40	45
307	His Val Ala Tyr Phe Val Pro Met Ser Lys Ile Ala Asp Phe Leu Lys		
308	50	55	60
310	Ala Gly Cys Glu Val Thr Ile Leu Phe Ala Asp Leu His Ala Tyr Leu		
311	65	70	75
313	Asp Asn Met Lys Ala Pro Trp Glu Leu Leu Glu Leu Arg Val Ser Tyr		80
314	85	90	95
316	Tyr Glu Asn Val Ile Lys Ala Met Leu Glu Ser Ile Gly Val Pro Leu		
317	100	105	110
319	Glu Lys Leu Lys Phe Ile Lys Gly Thr Asp Tyr Gln Leu Ser Lys Glu		
320	115	120	125
322	Tyr Thr Leu Asp Val Tyr Arg Leu Ser Ser Val Val Thr Gln His Asp		
323	130	135	140
325	Ser Lys Lys Ala Gly Ala Glu Val Val Lys Gln Val Glu His Pro Leu		
326	145	150	155
328	Leu Ser Gly Leu Leu Tyr Pro Gly Leu Gln Ala Leu Asp Glu Glu Tyr		160
329	165	170	175
331	Leu Lys Val Asp Ala Gln Phe Gly Gly Ile Asp Gln Arg Lys Ile Phe		
332	180	185	190
334	Thr Phe Ala Glu Lys Tyr Leu Pro Ala Leu Gly Tyr Ser Lys Arg Val		
335	195	200	205
337	His Leu Met Asn Pro Met Val Pro Gly Leu Thr Gly Ser Lys Met Ser		
338	210	215	220
340	Ser Ser Glu Glu Glu Ser Lys Ile Asp Leu Leu Asp Arg Lys Glu Asp		
341	225	230	235
343	Val Lys Lys Lys Leu Lys Lys Ala Phe Cys Glu Pro Gly Asn Val Glu		240
344	245	250	255
346	Asn Asn Gly Val Leu Ser Phe Ile Lys His Val Leu Phe Pro Leu Lys		
347	260	265	270
349	Ser Glu Phe Val Ile Leu Arg Asp Glu Lys Trp Gly Gly Asn Lys Thr		
350	275	280	285
352	Tyr Thr Ala Tyr Val Asp Leu Glu Lys Asp Phe Ala Ala Glu Val Val		
353	290	295	300
355	His Pro Gly Asp Leu Lys Asn Ser Val Glu Val Ala Leu Asn Lys Leu		
356	305	310	315
358	Leu Asp Pro Ile Arg Glu Lys Phe Asn Thr Pro Ala Leu Lys Lys Leu		320
359	325	330	335
361	Ala Ser Ala Ala Tyr Pro Asp Pro Ser Lys Gln Lys Pro Met Ala Lys		
362	340	345	350
364	Gly Pro Ala Lys Asn Ser Glu Pro Glu Glu Val Ile Pro Ser Arg Leu		
365	355	360	365
367	Asp Ile Arg Val Gly Lys Ile Ile Thr Val Glu Lys His Pro Asp Ala		
368	370	375	380
370	Asp Ser Leu Tyr Val Glu Lys Ile Asp Val Gly Glu Ala Glu Pro Arg		
371	385	390	395
373	Thr Val Val Ser Gly Leu Val Gln Phe Val Pro Lys Glu Glu Leu Gln		400
374	405	410	415
376	Asp Arg Leu Val Val Val Leu Cys Asn Leu Lys Pro Gln Lys Met Arg		
377	420	425	430

**VERIFICATION SUMMARY**

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Input Set : A:\Pto.da

Output Set: N:\CRF3\04112001\I813718.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/813,718

DATE: 04/05/2001

TIME: 08:08:35

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Output Set: N:\CRF3\04052001\I813718.raw

Does Not Comply  
Corrected Diskette Needed

3 <110> APPLICANT: Schimmel, Paul  
4 Wakasugi, Keisuke  
6 <120> TITLE OF INVENTION: Human Aminoacyl-tRNA Synthetase Polypeptides Useful For  
7 The Regulation of Angiogenesis  
9 <130> FILE REFERENCE: 00-221  
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/813,718  
C--> 12 <141> CURRENT FILING DATE: 2001-03-21  
14 <160> NUMBER OF SEQ ID NOS: 58  
16 <170> SOFTWARE: PatentIn Ver. 2.0

## ERRORED SEQUENCES

3151 <210> SEQ ID NO: 58  
3152 <211> LENGTH: 5  
3153 <212> TYPE: PRT  
3154 <213> ORGANISM: Homo sapiens  
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3157 Ser Ala Lys Gly Ile  
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**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/813,718

DATE: 04/05/2001

TIME: 08:08:36

Input Set : A:\tsri-8170.txt

Output Set: N:\CRF3\04052001\I813718.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:3163 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:58

M:332 Repeated in SeqNo=58